



C007-7022US0 - Sequence Listing
SEQUENCE LISTING

<110> Cohen, Edward H

<120> CONFORMATION SPECIFIC ANTIBODIES

<130> C007-7022US0

<140> US 10/589,956

<141> 2007-09-20

<160> 61

<170> PatentIn version 3.4

<210> 1

<211> 5

<212> PRT

<213> Homo sapiens

<400> 1

Arg Tyr Val Met Trp
1 5

<210> 2

<211> 17

<212> PRT

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Tyr Ile Trp Pro Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val Lys
1 5 10 15

Gly

<210> 3

<211> 11

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Ser Tyr Asp Phe Trp Ser Asn Ala Phe Asp Ile
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<221> misc

<222> (1)..(13)

C007-7022US0 - Sequence Listing

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Xaa Ser Xaa Asp Xaa Xaa Ser Xaa Ala Xaa Xaa Xaa Xaa
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<222> (1)..(11)

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Ser Tyr Asp Leu Trp Ser Asn Ala Tyr Asp Lys
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<222> (1)..(11)

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Ser Tyr Asp Leu Trp Ser Asn Ala Tyr Asp Lys
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Arg Ala Ser Gln Ser Ile Gly Ser Tyr Leu Asn
1 5 10

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Ala Ala Ser Ser Leu Gln Ser
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<213> Homo sapiens

<400> 9

Gln Gln Ser Tyr Ser Thr Pro Ser
1 5

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<400> 10

His Tyr Gly Met Ser
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<210> 11

<211> 17

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<400> 11

Val Ile Ser Pro Ser Gly Gly Arg Thr Leu Tyr Ala Asp Ser Val Lys
1 5 10 15

Gly

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His Tyr Ser Tyr Ala Met Asp Val
1 5

<210> 13

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<213> Homo sapiens

<400> 13

Thr Ala Ser Gln Ser Val Asp Ser Asn Leu Ala
1 5 10

<210> 14

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<213> Homo sapiens

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Gly Ala Ser Thr Arg Ala Thr
1 5

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Gln Gln Tyr Asn Lys Trp Pro Pro Tyr Ser
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His Tyr Ser Met Gln
1 5

<210> 17
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<213> Homo sapiens

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Tyr Ile Gly Ser Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val Lys
1 5 10 15

Gly

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Gly Thr Tyr Asn Thr Ser Pro Phe Asp Tyr
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<210> 19
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<400> 19

Ser Gly Asp Ala Leu Gly Gln Lys Tyr Ala Ser
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Gln Asp Ser Lys Arg Pro Ser
 1 5

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Gln Ala Trp Asp Thr Thr Ala Tyr Val
 1 5

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Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
 1 5 10 15

Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Gly Ser Tyr
 20 25 30

Leu Asn Trp Tyr Gln Gln Lys Thr Gly Lys Ala Pro Lys Ala Leu Ile
 35 40 45

Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly
 50 55 60

Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Leu
 65 70 75 80

Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Ser Thr Pro Ser
 85 90 95

Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr
 100 105

<210> 23
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 <213> Homo sapiens

<400> 23

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Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Arg Tyr
20 25 30

Val Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Tyr Ile Trp Pro Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Ser Ser Tyr Asp Phe Trp Ser Asn Ala Phe Asp Ile Trp Gly Gln
100 105 110

Gly Thr Met Val Thr Val Ser Ser
115 120

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<222> (1)..(108)

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Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
1 5 10 15

Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Gly Ser Tyr
20 25 30

Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Ala Leu Ile
35 40 45

Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly
50 55 60

Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
65 70 75 80

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Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Ser Thr Pro Ser
85 90 95

Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr
100 105

<210> 25
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<212> PRT
<213> Homo sapiens

<220>
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<222> (1)..(120)

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Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Arg Tyr
20 25 30

Val Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45
ø

Ser Tyr Ile Trp Pro Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Ser Ser Tyr Asp Phe Trp Ser Asn Ala Phe Asp Ile Trp Gly Gln
100 105 110

Gly Thr Met Val Thr Val Ser Ser
115 120

<210> 26
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<400> 26

Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Val Ser Pro Gly
1 5 10 15

Glu Arg Val Thr Leu Ser Cys Thr Ala Ser Gln Ser Val Asp Ser Asn
20 25 30

Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu Val
35 40 45

Tyr Gly Ala Ser Thr Arg Ala Thr Gly Val Pro Ala Arg Phe Ser Gly
50 55 60

Ser Gly Ser Gly Thr Ala Phe Thr Leu Thr Ile Asp Ser Leu Gln Ser
65 70 75 80

Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Tyr Asn Lys Trp Pro Pro
85 90 95

Tyr Ser Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg Thr
100 105 110

<210> 27

<211> 117

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<222> (1)..(117)

<400> 27

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser His Tyr
20 25 30

Gly Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Val Ile Ser Pro Ser Gly Gly Arg Thr Leu Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

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Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Lys His Tyr Ser Tyr Ala Met Asp Val Trp Gly Gln Gly Thr Thr
100 105 110

Val Thr Val Ser Ser
115

<210> 28
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<212> PRT
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<222> (1)..(105)

<400> 28

Ser Val Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln Thr
1 5 10 15

Ala Ser Val Thr Cys Ser Gly Asp Ala Leu Gly Gln Lys Tyr Ala Ser
20 25 30

Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Val Leu Val Ile Phe Gln
35 40 45

Asp Ser Lys Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser Asn
50 55 60

Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Val Asp
65 70 75 80

Glu Ala Asp Tyr Tyr Cys Gln Ala Trp Asp Thr Thr Ala Tyr Val Phe
85 90 95

Gly Thr Gly Thr Lys Val Thr Val Leu
100 105

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<400> 29

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Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gln Gln
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser His Tyr
20 25 30

Ser Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Tyr Ile Gly Ser Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Glu Gly Thr Tyr Asn Thr Ser Pro Phe Asp Tyr Trp Gly Gln Gly
100 105 110

Thr Leu Val Thr Val Ser Ser
115

<210> 30
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<400> 30

Met Lys Asp Ser Cys Ile Thr Val Met Ala Met Ala Leu Leu Ser Gly
1 5 10 15

Phe Phe Phe Phe Ala Pro Ala Ser Ser Tyr Asn Leu Asp Val Arg Gly
20 25 30

Ala Arg Ser Phe Ser Pro Pro Arg Ala Gly Arg His Phe Gly Tyr Arg
35 40 45

Val Leu Gln Val Gly Asn Gly Val Ile Val Gly Ala Pro Gly Glu Gly
50 55 60

Asn Ser Thr Gly Ser Leu Tyr Gln Cys Gln Ser Gly Thr Gly His Cys
65 70 75 80

Leu Pro Val Thr Leu Arg Gly Ser Asn Tyr Thr Ser Lys Tyr Leu Gly
85 90 95

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Met Thr Leu Ala Thr Asp Pro Thr Asp Gly Ser Ile Leu Ala Cys Asp
100 105 110

Pro Gly Leu Ser Arg Thr Cys Asp Gln Asn Thr Tyr Leu Ser Gly Leu
115 120 125

Cys Tyr Leu Phe Arg Gln Asn Leu Gln Gly Pro Met Leu Gln Gly Arg
130 135 140

Pro Gly Phe Gln Glu Cys Ile Lys Gly Asn Val Asp Leu Val Phe Leu
145 150 155 160

Phe Asp Gly Ser Met Ser Leu Gln Pro Asp Glu Phe Gln Lys Ile Leu
165 170 175

Asp Phe Met Lys Asp Val Met Lys Lys Leu Ser Asn Thr Ser Tyr Gln
180 185 190

Phe Ala Ala Val Gln Phe Ser Thr Ser Tyr Lys Thr Glu Phe Asp Phe
195 200 205

Ser Asp Tyr Val Lys Trp Lys Asp Pro Asp Ala Leu Leu Lys His Val
210 215 220

Lys His Met Leu Leu Leu Thr Asn Thr Phe Gly Ala Ile Asn Tyr Val
225 230 235 240

Ala Thr Glu Val Phe Arg Glu Glu Leu Gly Ala Arg Pro Asp Ala Thr
245 250 255

Lys Val Leu Ile Ile Ile Thr Asp Gly Glu Ala Thr Asp Ser Gly Asn
260 265 270

Ile Asp Ala Ala Lys Asp Ile Ile Arg Tyr Ile Ile Gly Ile Gly Lys
275 280 285

His Phe Gln Thr Lys Glu Ser Gln Glu Thr Leu His Lys Phe Ala Ser
290 295 300

Lys Pro Ala Ser Glu Phe Val Lys Ile Leu Asp Thr Phe Glu Lys Leu
305 310 315 320

Lys Asp Leu Phe Thr Glu Leu Gln Lys Lys Ile Tyr Val Ile Glu Gly
325 330 335

Thr Ser Lys Gln Asp Leu Thr Ser Phe Asn Met Glu Leu Ser Ser Ser
340 345 350

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Gly Ile Ser Ala Asp Leu Ser Arg Gly His Ala Val Val Gly Ala Val
355 360 365

Gly Ala Lys Asp Trp Ala Gly Gly Phe Leu Asp Leu Lys Ala Asp Leu
370 375 380

Gln Asp Asp Thr Phe Ile Gly Asn Glu Pro Leu Thr Pro Glu Val Arg
385 390 395 400

Ala Gly Tyr Leu Gly Tyr Thr Val Thr Trp Leu Pro Ser Arg Gln Lys
405 410 415

Thr Ser Leu Leu Ala Ser Gly Ala Pro Arg Tyr Gln His Met Gly Arg
420 425 430

Val Leu Leu Phe Gln Glu Pro Gln Gly Gly Gly His Trp Ser Gln Val
435 440 445

Gln Thr Ile His Gly Thr Gln Ile Gly Ser Tyr Phe Gly Gly Glu Leu
450 455 460

Cys Gly Val Asp Val Asp Gln Asp Gly Glu Thr Glu Leu Leu Leu Ile
465 470 475 480

Gly Ala Pro Leu Phe Tyr Gly Glu Gln Arg Gly Gly Arg Val Phe Ile
485 490 495

Tyr Gln Arg Arg Gln Leu Gly Phe Glu Glu Val Ser Glu Leu Gln Gly
500 505 510

Asp Pro Gly Tyr Pro Leu Gly Arg Phe Gly Glu Ala Ile Thr Ala Leu
515 520 525

Thr Asp Ile Asn Gly Asp Gly Leu Val Asp Val Ala Val Gly Ala Pro
530 535 540

Leu Glu Glu Gln Gly Ala Val Tyr Ile Phe Asn Gly Arg His Gly Gly
545 550 555 560

Leu Ser Pro Gln Pro Ser Gln Arg Ile Glu Gly Thr Gln Val Leu Ser
565 570 575

Gly Ile Gln Trp Phe Gly Arg Ser Ile His Gly Val Lys Asp Leu Glu
580 585 590

Gly Asp Gly Leu Ala Asp Val Ala Val Gly Ala Glu Ser Gln Met Ile

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595

600

605

Val Leu Ser Ser Arg Pro Val Val Asp Met Val Thr Leu Met Ser Phe
 610 615 620

Ser Pro Ala Glu Ile Pro Val His Glu Val Glu Cys Ser Tyr Ser Thr
 625 630 635 640

Ser Asn Lys Met Lys Glu Gly Val Asn Ile Thr Ile Cys Phe Gln Ile
 645 650 655

Lys Ser Leu Tyr Pro Gln Phe Gln Gly Arg Leu Val Ala Asn Leu Thr
 660 665 670

Tyr Thr Leu Gln Leu Asp Gly His Arg Thr Arg Arg Arg Gly Leu Phe
 675 680 685

Pro Gly Gly Arg His Glu Leu Arg Arg Asn Ile Ala Val Thr Thr Ser
 690 695 700

Met Ser Cys Thr Asp Phe Ser Phe His Phe Pro Val Cys Val Gln Asp
 705 710 715 720

Leu Ile Ser Pro Ile Asn Val Ser Leu Asn Phe Ser Leu Trp Glu Glu
 725 730 735

Glu Gly Thr Pro Arg Asp Gln Arg Ala Gln Gly Lys Asp Ile Pro Pro
 740 745 750

Leu Leu Arg Pro Ser Leu His Ser Glu Thr Trp Glu Ile Pro Phe Glu
 755 760 765

Lys Asn Cys Gly Glu Asp Lys Lys Cys Glu Ala Asn Leu Arg Val Ser
 770 775 780

Phe Ser Pro Ala Arg Ser Arg Ala Leu Arg Leu Thr Ala Phe Ala Ser
 785 790 795 800

Leu Ser Val Glu Leu Ser Leu Ser Asn Leu Glu Glu Asp Ala Tyr Trp
 805 810 815

Val Gln Leu Asp Leu His Phe Pro Pro Gly Leu Ser Phe Arg Lys Val
 820 825 830

Glu Met Leu Lys Pro His Ser Gln Ile Pro Val Ser Cys Glu Glu Leu
 835 840 845

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Pro Glu Glu Ser Arg Leu Leu Ser Arg Ala Leu Ser Cys Asn Val Ser
850 855 860

Ser Pro Ile Phe Lys Ala Gly His Ser Val Ala Leu Gln Met Met Phe
865 870 875 880

Asn Thr Leu Val Asn Ser Ser Trp Gly Asp Ser Val Glu Leu His Ala
885 890 895

Asn Val Thr Cys Asn Asn Glu Asp Ser Asp Leu Leu Glu Asp Asn Ser
900 905 910

Ala Thr Thr Ile Ile Pro Ile Leu Tyr Pro Ile Asn Ile Leu Ile Gln
915 920 925

Asp Gln Glu Asp Ser Thr Leu Tyr Val Ser Phe Thr Pro Lys Gly Pro
930 935 940

Lys Ile His Gln Val Lys His Met Tyr Gln Val Arg Ile Gln Pro Ser
945 950 955 960

Ile His Asp His Asn Ile Pro Thr Leu Glu Ala Val Val Gly Val Pro
965 970 975

Gln Pro Pro Ser Glu Gly Pro Ile Thr His Gln Trp Ser Val Gln Met
980 985 990

Glu Pro Pro Val Pro Cys His Tyr Glu Asp Leu Glu Arg Leu Pro Asp
995 1000 1005

Ala Ala Glu Pro Cys Leu Pro Gly Ala Leu Phe Arg Cys Pro Val
1010 1015 1020

Val Phe Arg Gln Glu Ile Leu Val Gln Val Ile Gly Thr Leu Glu
1025 1030 1035

Leu Val Gly Glu Ile Glu Ala Ser Ser Met Phe Ser Leu Cys Ser
1040 1045 1050

Ser Leu Ser Ile Ser Phe Asn Ser Ser Lys His Phe His Leu Tyr
1055 1060 1065

Gly Ser Asn Ala Ser Leu Ala Gln Val Val Met Lys Val Asp Val
1070 1075 1080

Val Tyr Glu Lys Gln Met Leu Tyr Leu Tyr Val Leu Ser Gly Ile
1085 1090 1095

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Gly Gly Leu Leu Leu Leu Leu Leu Ile Phe Ile Val Leu Tyr Lys
1100 1105 1110

Val Gly Phe Phe Lys Arg Asn Leu Lys Glu Lys Met Glu Ala Gly
1115 1120 1125

Arg Gly Val Pro Asn Gly Ile Pro Ala Glu Asp Ser Glu Gln Leu
1130 1135 1140

Ala Ser Gly Gln Glu Ala Gly Asp Pro Gly Cys Leu Lys Pro Leu
1145 1150 1155

His Glu Lys Asp Ser Glu Ser Gly Gly Gly Lys Asp
1160 1165 1170

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<213> Homo sapiens

<400> 31

Met Leu Gly Leu Arg Pro Pro Leu Leu Ala Leu Val Gly Leu Leu Ser
1 5 10 15

Leu Gly Cys Val Leu Ser Gln Glu Cys Thr Lys Phe Lys Val Ser Ser
20 25 30

Cys Arg Glu Cys Ile Glu Ser Gly Pro Gly Cys Thr Trp Cys Gln Lys
35 40 45

Leu Asn Phe Thr Gly Pro Gly Asp Pro Asp Ser Ile Arg Cys Asp Thr
50 55 60

Arg Pro Gln Leu Leu Met Arg Gly Cys Ala Ala Asp Asp Ile Met Asp
65 70 75 80

Pro Thr Ser Leu Ala Glu Thr Gln Glu Asp His Asn Gly Gly Gln Lys
85 90 95

Gln Leu Ser Pro Gln Lys Val Thr Leu Tyr Leu Arg Pro Gly Gln Ala
100 105 110

Ala Ala Phe Asn Val Thr Phe Arg Arg Ala Lys Gly Tyr Pro Ile Asp
115 120 125

Leu Tyr Tyr Leu Met Asp Leu Ser Tyr Ser Met Leu Asp Asp Leu Arg
130 135 140

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Asn Val Lys Lys Leu Gly Gly Asp Leu Leu Arg Ala Leu Asn Glu Leu
 145 150 155 160
 Thr Glu Ser Gly Arg Ile Gly Phe Gly Ser Phe Val Asp Lys Thr Val
 165 170 175
 Leu Pro Phe Val Asn Thr Ile Ile Pro Asp Lys Leu Arg Asn Pro Cys
 180 185 190
 Pro Asn Lys Glu Lys Glu Cys Gln Pro Pro Phe Ala Phe Arg His Val
 195 200 205
 Leu Lys Leu Thr Asn Asn Ser Asn Gln Phe Gln Thr Glu Val Gly Lys
 210 215 220
 Gln Leu Ile Ser Gly Asn Leu Asp Ala Pro Glu Gly Gly Leu Asp Ala
 225 230 235 240
 Met Met Gln Val Ala Ala Cys Pro Glu Glu Ile Gly Trp Arg Asn Val
 245 250 255
 Thr Arg Leu Leu Val Phe Ala Thr Asp Asp Gly Phe His Phe Ala Gly
 260 265 270
 Asp Gly Lys Leu Gly Ala Ile Leu Thr Pro Asn Asp Gly Arg Cys His
 275 280 285
 Leu Glu Asp Asn Leu Tyr Lys Arg Ser Asn Glu Phe Asp Val Pro Ser
 290 295 300
 Val Gly Gln Leu Ala His Lys Leu Ala Glu Asn Asn Ile Gln Pro Ile
 305 310 315 320
 Phe Ala Val Thr Ser Arg Met Val Lys Thr Tyr Glu Lys Leu Thr Glu
 325 330 335
 Ile Ile Pro Lys Ser Ala Val Gly Glu Leu Ser Glu Asp Ser Ser Asn
 340 345 350
 Val Val His Leu Ile Lys Asn Ala Tyr Asn Lys Leu Ser Ser Arg Val
 355 360 365
 Phe Leu Asp His Asn Ala Leu Pro Asp Thr Leu Lys Val Thr Tyr Asp
 370 375 380
 Ser Phe Cys Ser Asn Gly Val Thr His Arg Asn Gln Pro Arg Gly Asp
 385 390 395 400

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Cys Asp Gly Val Gln Ile Asn Val Pro Ile Thr Phe Gln Val Lys Val
 405 410 415
 Thr Ala Thr Glu Cys Ile Gln Glu Gln Ser Phe Val Ile Arg Ala Leu
 420 425 430
 Gly Phe Thr Asp Ile Val Thr Val Gln Val Leu Pro Gln Cys Glu Cys
 435 440 445
 Arg Cys Arg Asp Gln Ser Arg Asp Arg Ser Leu Cys His Gly Lys Gly
 450 455 460
 Phe Leu Glu Cys Gly Ile Cys Arg Cys Asp Thr Gly Tyr Ile Gly Lys
 465 470 475 480
 Asn Cys Glu Cys Gln Thr Gln Gly Arg Ser Ser Gln Glu Leu Glu Gly
 485 490 495
 Ser Cys Arg Lys Asp Asn Asn Ser Ile Ile Cys Ser Gly Leu Gly Asp
 500 505 510
 Cys Val Cys Gly Gln Cys Leu Cys His Thr Ser Asp Val Pro Gly Lys
 515 520 525
 Leu Ile Tyr Gly Gln Tyr Cys Glu Cys Asp Thr Ile Asn Cys Glu Arg
 530 535 540
 Tyr Asn Gly Gln Val Cys Gly Gly Pro Gly Arg Gly Leu Cys Phe Cys
 545 550 555 560
 Gly Lys Cys Arg Cys His Pro Gly Phe Glu Gly Ser Ala Cys Gln Cys
 565 570 575
 Glu Arg Thr Thr Glu Gly Cys Leu Asn Pro Arg Arg Val Glu Cys Ser
 580 585 590
 Gly Arg Gly Arg Cys Arg Cys Asn Val Cys Glu Cys His Ser Gly Tyr
 595 600 605
 Gln Leu Pro Leu Cys Gln Glu Cys Pro Gly Cys Pro Ser Pro Cys Gly
 610 615 620
 Lys Tyr Ile Ser Cys Ala Glu Cys Leu Lys Phe Glu Lys Gly Pro Phe
 625 630 635 640

Gly Lys Asn Cys Ser Ala Ala Cys Pro Gly Leu Gln Leu Ser Asn Asn
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645

650

655

Pro Val Lys Gly Arg Thr Cys Lys Glu Arg Asp Ser Glu Gly Cys Trp
660 665 670

Val Ala Tyr Thr Leu Glu Gln Gln Asp Gly Met Asp Arg Tyr Leu Ile
675 680 685

Tyr Val Asp Glu Ser Arg Glu Cys Val Ala Gly Pro Asn Ile Ala Ala
690 695 700

Asn Gly Gly Thr Val Ala Gly Ile Val Leu Ile Gly Ile Leu Leu Leu
705 710 715 720

Val Ile Trp Lys Ala Leu Ile His Leu Ser Asp Leu Arg Glu Tyr Arg
725 730 735

Arg Phe Glu Lys Glu Lys Leu Lys Ser Gln Trp Asn Asn Asp Asn Pro
740 745 750

Leu Phe Lys Ser Ala Thr Thr Thr Val Met Asn Pro Lys Phe Ala Glu
755 760 765

Ser

<210> 32
<211> 323
<212> PRT
<213> Homo sapiens

<400> 32

Gly Asn Val Asp Leu Val Phe Leu Phe Asp Gly Ser Met Ser Leu Gln
1 5 10 15

Pro Asp Glu Phe Gln Lys Ile Leu Asp Phe Met Lys Asp Val Met Lys
20 25 30

Lys Leu Ser Asn Thr Ser Tyr Gln Phe Ala Ala Val Gln Phe Ser Thr
35 40 45

Ser Glu Glu Glu Glu Glu Glu Glu Asx Thr Thr Ser His His His
50 55 60

His His His His His His His His His His His His His His Thr Thr
65 70 75 80

Thr ser ser Glu Glu Glu Glu Glu Glu Glu Glu Ser Ser Ser Tyr Lys

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85

90

95

Thr Glu Phe Asp Phe Ser Asp Tyr Val Lys Arg Lys Asp Pro Asp Ala
100 105 110

Leu Leu Lys His Val Lys His Met Leu Leu Leu Thr Asn Thr Phe Gly
115 120 125

Ala Ile Asn Tyr Val Ala Thr Glu Val Phe Arg Glu Glu Leu Gly Ala
130 135 140

Glu Glu Glu Ser Asx His His His His His His His Thr Thr His His
145 150 155 160

His His Thr Ser Ser Asx His His His His His His His His His His
165 170 175

His Thr Thr Thr Gly Gly Gly Thr Arg Pro Asp Ala Thr Lys Val Leu
180 185 190

Ile Ile Ile Thr Asp Gly Glu Ala Thr Asp Ser Gly Asn Ile Asp Ala
195 200 205

Ala Lys Asp Ile Ile Arg Tyr Ile Ile Gly Ile Gly Lys His Phe Gln
210 215 220

Thr Lys Glu Ser Gln Glu Thr Leu His Lys Thr Thr Ser Glu Glu Glu
225 230 235 240

Glu Glu Glu Glu Glu Ser Ser Gly Gly Gly Thr Thr Ser Glu Glu Glu
245 250 255

Glu Glu Glu Ser Ser Ser Thr Thr Thr Gly Gly Gly Gly Thr Thr Phe
260 265 270

Ala Ser Lys Pro Ala Ser Glu Phe Val Lys Ile Leu Asp Thr Phe Glu
275 280 285

Lys Leu Lys Asp Leu Phe Thr Glu Leu Gln Lys Lys Ile Thr Ser Ser
290 295 300

Ser His His His His Glu Glu Glu Thr Thr Thr Thr Thr Thr Thr
305 310 315 320

Thr Thr Thr

<210> 33

C007-7022US0 - Sequence Listing

<211> 109
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<213> Homo sapiens

<220>
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<222> (1)..(109)

<400> 33

Gln Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val
1 5 10 15

Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Gly Ser
20 25 30

Tyr Leu Asn Trp Tyr Gln Gln Lys Thr Gly Lys Ala Pro Lys Ala Leu
35 40 45

Ile Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser
50 55 60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65 70 75 80

Leu Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Ser Thr Pro
85 90 95

Ser Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr
100 105

<210> 34
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<400> 34

Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Val Ser Pro
1 5 10 15

Gly Glu Arg Val Thr Leu Ser Cys Thr Ala Ser Gln Ser Val Asp Ser
20 25 30

Asn Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu
35 40 45

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Val Tyr Gly Ala Ser Thr Arg Ala Thr Gly Val Pro Ala Arg Phe Ser
50 55 60

Gly Ser Gly Ser Gly Thr Ala Phe Thr Leu Tyr Ile Asp Ser Leu Gln
65 70 75 80

Ser Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Tyr Asn Lys Trp Pro
85 90 95

Pro Tyr Ser Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg Thr
100 105 110

<210> 35
<211> 106
<212> PRT
<213> Homo sapiens

<220>
<221> Protein
<222> (1)..(106)

<400> 35

Gln Ser Val Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
1 5 10 15

Thr Ala Ser Val Thr Cys Ser Gly Asp Ala Leu Gly Gln Lys Tyr Ala
20 25 30

Ser Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Val Leu Val Ile Phe
35 40 45

Gln Asp Ser Lys Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
50 55 60

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Val
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln Ala Trp Asp Thr Thr Ala Tyr Val
85 90 95

Phe Gly Thr Gly Thr Lys Val Thr Val Leu
100 105

<210> 36
<211> 120
<212> PRT
<213> Homo sapiens

<220>

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<221> Protein
<222> (1)..(120)

<400> 36

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Arg Tyr
20 25 30

Val Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Tyr Ile Trp Pro Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Ser Ser Tyr Asp Phe Trp Ser Asn Ala Phe Asp Ile Trp Gly Gln
100 105 110

Gly Thr Met Val Thr Val Ser Ser
115 120

<210> 37
<211> 117
<212> PRT
<213> Homo sapiens

<220>
<221> Protein
<222> (1)..(117)

<400> 37

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser His Tyr
20 25 30

Gly Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Val Ile Ser Pro Ser Gly Gly Arg Thr Leu Tyr Ala Asp Ser Val
50 55 60

C007-7022US0 - Sequence Listing

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Lys His Tyr Ser Tyr Ala Met Asp Val Trp Gly Gln Gly Thr Thr
100 105 110

Val Thr Val Ser Ser
115

<210> 38
<211> 122
<212> PRT
<213> Homo sapiens

<220>
<221> misc_feature
<222> (100)..(100)
<223> Xaa can be any naturally occurring amino acid

<400> 38

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser His Tyr
20 25 30

Ser Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Tyr Ile Gly Ser Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Arg Gly Xaa Ala Ala Thr Tyr Asn Thr Ser Pro Phe Asp Tyr Trp
100 105 110

Gly Gln Gly Thr Leu Val Thr Val Ser Ser
115 120

C007-7022US0 - Sequence Listing

<210> 39
<211> 321
<212> DNA
<213> Homo sapiens

<220>
<221> DNA
<222> (1)..(321)

<400> 39
caagacatcc agatgaccca gtctccatcc tccctgtctg catctgtagg agacagagtc 60
accatcactt gccgggcaag tcagagcatt ggcagctact taaactggta tcagcagaaa 120
acaggggaaag cccctaaggc cctgatctat gctgcatcca gtttgcaaag tgggggtccca 180
tcaaggttca gtggcagtgg gtctgggaca gatttcactc tcaccatcag tagtctgcaa 240
cttgaagatt ttgcaactta ctactgtcaa cagagttaca gtaccccctc gttcggccaa 300
gggaccaagg tggaaatcaa a 321

<210> 40
<211> 327
<212> DNA
<213> Homo sapiens

<400> 40
caagacatcc agatgaccca gtctccagcc accctgtctg tgtctccagg ggaaagagtc 60
accctctcct gcacggccag tcagagtgtt gacagcaact tagcctggta tcagcaaaaa 120
cctggccagg ctcccagact cctcgtctat ggtgcatcca ctagggccac tgggtgtccca 180
gccagggttca gtggcagtgg gtctgggaca gcgttcactc tcaccatcga cagcctgcag 240
tctgaagatt ttgcagttta ttactgtcag cagtataata agtggcctcc gtactccttt 300
ggccagggga ccaagctgga gatcaag 327

<210> 41
<211> 318
<212> DNA
<213> Homo sapiens

<220>
<221> DNA
<222> (1)..(318)

<400> 41
cagagcgtct tgactcagcc accctcagtg tccgtctccc caggacagac agccagcgtc 60
acttgctctg gagatgcatt gggacaaaaa tatgcttcct ggtatcaaca gaagccaggc 120
cagtccccctg tactgggtcat ctttcaagat tccaagcggc cctcagggat ccctgagcgg 180
ttctctggct ccaattctgg gaacacagcc actctgacca tcagcgggac ccaggctgtg 240
gatgaggccg actattattg tcaggcgtgg gacactacag cttatgtctt cggaactggg 300

C007-7022US0 - Sequence Listing

accaaggtca ccgtccta

318

<210> 42
 <211> 360
 <212> DNA
 <213> Homo sapiens

<220>
 <221> DNA
 <222> (1)..(360)

<400> 42
 gaagttcaat tgtagagtc tggtagcggt cttgttcagc ctggtgggtc ttacgtctt 60
 tcttgcgctg cttccggatt cactttctct cgttacgtta tgtggtgggt tcgccaagct 120
 cctggtaaag gtttggagtg ggtttcttat atctggcctt ctggtggcaa tacttattat 180
 gctgactccg ttaaagggtcg cttcactatc tctagagaca actctaagaa tactctctac 240
 ttgcagatga acagcttaag ggctgaggac actgcagtct actattgtgc gagtagctac 300
 gatttttggg gtaatgcttt tgatatctgg ggccaaggga caatggtcac cgtctcaagc 360

<210> 43
 <211> 351
 <212> DNA
 <213> Homo sapiens

<400> 43
 gaagttcaat tgtagagtc tggtagcggt cttgttcagc ctggtgggtc ttacgtctt 60
 tcttgcgctg cttccggatt cactttctct cattacggtg tgtcttgggt tcgccaagct 120
 cctggtaaag gtttggagtg ggtttctgtt atctctcctt ctggtggccg tactctttat 180
 gctgactccg ttaaagggtcg cttcactatc tctagagaca actctaagaa tactctctac 240
 ttgcagatga acagcttaag ggctgaggac actgcagtct actattgtgc gaaacattac 300
 tcctacgcta tggacgtctg gggccaaggg accacggtca ccgtctcaag c 351

<210> 44
 <211> 357
 <212> DNA
 <213> Homo sapiens

<400> 44
 gaagttcaat tgtagagtc tggtagcggt cttgttcagc ctggtgggtc ttacgtctt 60
 tcttgcgctg cttccggatt cactttctct cattactcta tgcagtgggt tcgccaagct 120
 cctggtaaag gtttggagtg ggtttcttat atcggttctt ctggtggcaa tacttattat 180
 gctgactccg ttaaagggtcg cttcactatc tctagagaca actctaagaa tactctctac 240
 ttgcagatga acagcttaag ggctgaggac actgcagtct actattgtgc gagagggacc 300

tataacacct cccctttga ctactggggc caggaaccc tggcaccgt ctcaagc

357

<210> 45
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 45

Ala Glu Tyr Phe Gln His Trp Gly Gln Gly Thr Leu Val Thr Val Ser
 1 5 10 15

Ser

<210> 46
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 46

Tyr Trp Tyr Phe Asp Leu Trp Gly Arg Gly Thr Leu Val Thr Val Ser
 1 5 10 15

Ser

<210> 47
 <211> 15
 <212> PRT
 <213> Homo sapiens

<400> 47

Ala Phe Asp Ile Trp Gly Gln Gly Thr Met Val Thr Val Ser Ser
 1 5 10 15

<210> 48
 <211> 29
 <212> PRT
 <213> Homo sapiens

<400> 48

Tyr Phe Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Phe
 1 5 10 15

Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
 20 25

<210> 49
 <211> 16
 <212> PRT
 <213> Homo sapiens

C007-7022US0 - Sequence Listing

<400> 49

Asn Trp Phe Asp Pro Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
1 5 10 15

<210> 50

<211> 20

<212> PRT

<213> Homo sapiens

<400> 50

Tyr Tyr Tyr Tyr Tyr Gly Met Asp Val Trp Gly Gln Gly Thr Thr Val
1 5 10 15

Thr Val Ser Ser
20

<210> 51

<211> 120

<212> PRT

<213> Homo sapiens

<400> 51

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Arg Tyr
20 25 30

Val Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Tyr Ile Trp Pro Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Ser Ser Tyr Asp Tyr Trp Ser Asn Ala Phe Asp Ser Trp Gly Gln
100 105 110

Gly Thr Met Val Thr Val Ser Ser
115 120

<210> 52

<211> 120

C007-7022US0 - Sequence Listing

<212> PRT
<213> Homo sapiens

<400> 52

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Arg Tyr
20 25 30

Val Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Tyr Ile Trp Pro Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Ser Ser Phe Asp Phe Trp Ser Asn Ala Phe Asp Met Trp Gly Gln
100 105 110

Gly Thr Met Val Thr Val Ser Ser
115 120

<210> 53
<211> 120
<212> PRT
<213> Homo sapiens

<400> 53

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Arg Tyr
20 25 30

Val Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Tyr Ile Trp Pro Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Leu Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

C007-7022US0 - Sequence Listing

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Ser Ser Tyr Asp Leu Trp Ser Asn Ala Phe Asp Lys Trp Gly Gln
100 105 110

Gly Thr Met Val Thr Val Ser Ser
115 120

<210> 54
<211> 120
<212> PRT
<213> Homo sapiens

<400> 54

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Arg Tyr
20 25 30

Val Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Tyr Ile Trp Pro Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Leu Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Asn Ser Tyr Asp Phe Arg Ser Asn Ala Phe Ala Val Trp Gly Gln
100 105 110

Gly Thr Met Val Thr Val Ser Ser
115 120

<210> 55
<211> 120
<212> PRT
<213> Homo sapiens

<220>
<221> Protein
<222> (1)..(120)

<400> 55

c007-7022US0 - Sequence Listing

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Arg Tyr
20 25 30

Val Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Tyr Ile Trp Pro Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Ser Ser Tyr Asp Leu Trp Ser Tyr Ala Phe Glu Ile Trp Gly Gln
100 105 110

Gly Thr Met Val Thr Val Ser Ser
115 120

<210> 56
<211> 120
<212> PRT
<213> Homo sapiens

<220>
<221> Protein
<222> (1)..(120)

<400> 56

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Arg Tyr
20 25 30

Val Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Tyr Ile Trp Pro Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

c007-7022US0 - Sequence Listing

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Ser Ser Tyr Asp Phe Trp Ser Asn Ala Tyr Ala Asn Trp Gly Gln
100 105 110

Gly Thr Met Val Thr Val Ser Ser
115 120

<210> 57
<211> 120
<212> PRT
<213> Homo sapiens

<400> 57

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Arg Tyr
20 25 30

Val Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Tyr Ile Trp Pro Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Leu Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Asn Ser Phe Asp Phe Trp Ser Asn Ala Phe Glu Leu Trp Gly Gln
100 105 110

Gly Thr Met Val Thr Val Ser Ser
115 120

<210> 58
<211> 639
<212> DNA
<213> Homo sapiens

<400> 58

gacatccaga tgaccagtc tccatcctcc ctgtctgcat ctgtaggaga cagagtcacc 60

atcacttgcc gggcaagtca gagcattggc agctacttaa actggtatca gcagaaacca 120

gggaaagccc ctaaggccct gatctatgct gcatccagtt tgcaaagtgg ggtcccatca 180

C007-7022US0 - Sequence Listing

aggttcagtg gcagtggggtc tgggacagat ttcactctca ccatcagtag tctgcaacct	240
gaagatttttg caacttacta ctgtcaacag agttacagta cccctcgtt cggccaaggg	300
accaaggttg aaatcaaaag aactgtggct gcaccatctg tcttcatctt cccgccatct	360
gatgagcagt tgaaatcttg aactgcctct gttgtgtgcc tgctgaataa cttctatccc	420
agagaggcca aagtacagtg gaaggtggat aacgccctcc aatcgggtaa ctcccaggag	480
agtgtcacag agcaggacag caaggacagc acctacagcc tcagcagcac cctgacgctg	540
agcaaagcag actacgagaa acacaaagtc tacgcctgcg aagtcacca tcagggcctg	600
agctcgcccc tcacaaagag cttcaacagg ggagagtgt	639

<210> 59
 <211> 1344
 <212> DNA
 <213> Homo sapiens

<400> 59	
gaagttcaat tgtagagtc tgggtggcggc cttgttcagc ctggtgggtc ttacgtctt	60
tcttgcgctg cttccggatt cactttctct cgttacgta tgtggtgggt tcgccaagct	120
cctggtaaaag gtttggagtg ggtttcttat atctggcctt ctggtggcaa tacttattat	180
gctgactccg ttaaaggctc cttcactatc tctagagaca actctaagaa tactctctac	240
ttgcagatga acagcttaag ggctgaggac actgcagtct actattgtgc gagtagctac	300
gatttttggg gtaatgcttt tgatatctgg ggccaagggg caatggtcac cgtctcaagc	360
gcctccacca agggcccatc ggtcttccc ctagcgccct gctccaggag cacctccgag	420
agcacagccg ccctgggctg cctgggtcaag gactacttcc ccgaaccggt gacggtgtcg	480
tggaactcag gcgccctgac cagcggcgct cacaccttcc cggctgtcct acagtctctc	540
ggactctact ccctcagcag cgtagtgacc gtgccctcca gcagcttggg cacgaagacc	600
tacacctgca acgtagatca caagcccagc aacaccaagg tggacaagag agttgagtcc	660
aaatatggtc ccccatgccc atcatgccc gcacctgagt tcctgggggg accatcagtc	720
ttcctgttcc ccccaaaacc caaggacact ctcagtatct cccggacccc tgaggtcacg	780
tgcgtgggtg tggacgtgag ccaggaagac cccgaggtcc agttcaactg gtacgtggat	840
ggcgtggagg tgcataatgc caagacaaag ccgcgggagg agcagttcaa cagcacgtac	900
cgtgtgggtc gcgtcctcac cgtcctgcac caggactggc tgaacggcaa ggagtacaag	960
tgcaagggtct ccaacaaagg cctcccgtcc tccatcgaga aaacctctc caaagccaaa	1020
gggcagcccc gagagccaca ggtgtacacc ctgcccccat cccaggagga gatgaccaag	1080
aaccagggtc gcctgacctg cctgggtcaaa ggcttctacc ccagcgacat cgccgtggag	1140
tgggagagca atgggcagcc ggagaacaac tacaagacca cgcctcccgt gctggactcc	1200

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gacggctcct tcttcctcta cagcaggcta accgtggaca agagcaggtg gcaggagggg 1260
aatgtcttct catgctccgt gatgcatgag gctctgcaca accactacac acagaagagc 1320
ctctccctgt ctctgggtaa atga 1344

<210> 60
<211> 213
<212> PRT
<213> Homo sapiens

<220>
<221> Protein
<222> (1)..(213)

<400> 60

Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
1 5 10 15

Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Gly Ser Tyr
20 25 30

Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Ala Leu Ile
35 40 45

Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly
50 55 60

Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
65 70 75 80

Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Ser Thr Pro Ser
85 90 95

Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala Ala Pro
100 105 110

Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser Gly Thr
115 120 125

Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu Ala Lys
130 135 140

Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly Asn Ser Gln Glu
145 150 155 160

Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr Ser Leu Ser Ser
165 170 175

c007-7022US0 - Sequence Listing

Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His Lys Val Tyr Ala
180 185 190

Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val Thr Lys Ser Phe
195 200 205

Asn Arg Gly Glu Cys
210

<210> 61
<211> 448
<212> PRT
<213> Homo sapiens

<220>
<221> Protein
<222> (1)..(448)

<400> 61

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Arg Tyr
20 25 30

Val Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Tyr Ile Trp Pro Ser Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Ser Ser Tyr Asp Phe Trp Ser Asn Ala Phe Asp Ile Trp Gly Gln
100 105 110

Gly Thr Met Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val
115 120 125

Phe Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala
130 135 140

Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser
145 150 155 160

C007-7022US0 - Sequence Listing

Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val
165 170 175

Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro
180 185 190

Ser Ser Ser Leu Gly Thr Lys Thr Tyr Thr Cys Asn Val Asp His Lys
195 200 205

Pro Ser Asn Thr Lys Val Asp Lys Arg Val Glu Ser Lys Tyr Gly Pro
210 215 220

Pro Cys Pro Ser Cys Pro Ala Pro Glu Phe Leu Gly Gly Pro Ser Val
225 230 235 240

Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr
245 250 255

Pro Glu Val Thr Cys Val Val Val Asp Val Ser Gln Glu Asp Pro Glu
260 265 270

Val Gln Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys
275 280 285

Thr Lys Pro Arg Glu Glu Gln Phe Asn Ser Thr Tyr Arg Val Val Ser
290 295 300

Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys
305 310 315 320

Cys Lys Val Ser Asn Lys Gly Leu Pro Ser Ser Ile Glu Lys Thr Ile
325 330 335

Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro
340 345 350

Pro Ser Gln Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu
355 360 365

Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Ser
370 375 380

Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp
385 390 395 400

Ser Asp Gly Ser Phe Phe Leu Tyr Ser Arg Leu Thr Val Asp Lys Ser

C007-7022US0 - Sequence Listing

405

410

415

Arg Trp Gln Glu Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala
420 425 430

Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Leu Gly Lys
435 440 445